

# Blast 2 Sequences results

9/600 493

PubMed

Entrez

BLAST

OMIM

Taxonomy

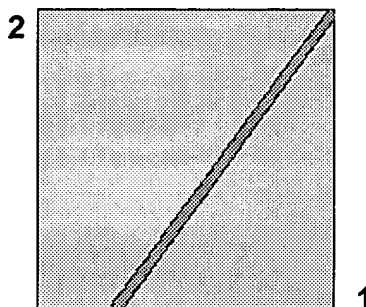
Structure

## BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.1.2 [Oct-19-2000]

Match:  Mismatch:  gap open:  gap extension: x\_dropoff:  expect:  wordsize:  ☐ Filter ☒ Align

**Sequence 1** gi\_59478 Hepatitis C virus core, E1, NS1/E2, NS2, NS3, NS4a, NS4b and NS5 genes **Length** 9408 (1 .. 9408)

**Sequence 2** gi\_451206 Hepatitis C virus gene for polyprotein precursor (NS2-NS3-NS4A-NS4B-NS5A-NS5B), complete cds. **Length** 6935 (1 .. 6935)



NOTE: The statistics (bitscore and expect value) is calculated based on the size of nr database

Score = 9508 bits (4945), Expect = 0.0

Identities = 6265/6887 (90%)

Strand = Plus / Plus

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[illegible]

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X2: 26 (50.0 bits)  
S1: 12 (23.8 bits)  
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